

09283034-021501

GAATTCGGCACGAGGTTTTTTTTTTTTTCCCTCTTTCTTTCTTTCTTTGCC
1 -----+-----+-----+-----+-----+ 60

ATCCGAAAGAGCTGTCAGCCGCCCGCGGGCTGCACCTAAAGGCGTCGGTAGGGGGATAAC
61 -----+-----+-----+-----+-----+ 120

AGTCAGAGACCCCTCCTGAAAGCAGGAGACGGGACGGTACCCCTCCGGCTCTGCGGGGCGG
121 -----+-----+-----+-----+-----+ 180

CTGCGGGCCCTCCGTTCTTTCCCCCTCCCCGAGAGACACTCTTCCTTTCCCCCACAAG
181 -----+-----+-----+-----+-----+ 240

ACACAGGGGCAGGAACGCGAGCGCTGCCCCCTCGCCATGGGAGGCGCTTCCTGCTGACG
241 -----+-----+-----+-----+-----+ 300

FIG. 1A

CTCGCCCTCCTCTCGGGCGCTGCTGTGCCGCTGCCAGGTTGACGGCTCCGGGGTGTTCGAG
301 -----+-----+-----+-----+-----+ 360

CTGAAGCTGCAGGAGTTTGTCAACAAGAAGGGGCTGCTCAGCAACCGCAACTGCTGCCGG
361 -----+-----+-----+-----+-----+ 420

GGGGGCGGCCCCGGAGGGCGCCGGGCAGCAGCAGTGCAGCTGCAAGACCTTCTTCCGCGTC
421 -----+-----+-----+-----+-----+ 480

TGCCTGPAGCACTACCAGGCCAGCGTCTCCCCGAGCCGCCCTGCACCTACGGCAGCGCC
481 -----+-----+-----+-----+-----+ 540

ATCACCCCCGTCTCGGGCCCACTCCTTCAACGTCCTCCGACGGCGGGCGGGCGCCGAC
541 -----+-----+-----+-----+-----+ 600

CCCGCCTTCAGCAACCCCATCCGCTTCCCCCTTCGGCTTCACCTGGCCCCGGCACCTTCTCG
601 -----+-----+-----+-----+-----+ 660

CTCATCATCGAGGCTCTGCACACCGACTCCCCCGACGACCTCACCACAGAAAACCCCGAG
661 -----+-----+-----+-----+-----+ 720

FIG. 1A (cont'd)

721 CGCCTCATCAGCCGCCTGGCCACCCAGAGGCACCTGGCGGTGGGCGAGGAGTGGTCCCAG 780

781 GACCTGCACAGCAGCGGCCG/ACCGACCTCAAGTACTCCTATCGCTTTGTGTGTGATGAG 840

841 CACTACTACGGGAAGGCTGCTCTGTCTTCGCCGGCCCCGTGACGACCGCTTCGGTCAC 900

901 TTCACCTGTGGAGAGCGTGGCGAGAAGGTCTGCAACCCAGGCTGGAAGGGCCAGTACTGC 960

961 ACTGAGCCGATTGCTTGCCCTGGGTGTGACGAGCAGCAGGCTTCTGCGACAAACCTGGG 1020

1021 GAATGCAAGTGCAGAGTGGGTGGCAGGGCGGTACTGTGACGAGTGCATCCGATACCCA 1080

1081 GGCTGCCCTGCACGGTACCTGTCAGCAGCCA/TGGCAGTGCAACTGCCAGGAAGGCTGGGGC 1140

FIG. 1A (cont'd)

[illegible]

FIG. 1A (cont'd)

1561 GACCTGGGGAACCTCCTACATATGCCAGTGCCAGGCTGGCTTCACTGGCAGGCACTGTGAC 1620

1621 GACAACGTGGACGATTGCGCCTCCTTCCCTGCGTCAATGGAGGGACCTGTCAGGATGGG 1680

1681 GTCAACGACTACTCCTGCACCTGCCCCCGGGATACAACGGGAAGAACTGCAGCAGCCG 1740

1741 GTGAGCAGATGCGAGCACAACCCCTGCCACAATGGGGCCACCTGCCACGAGAGAAGCAAC 1800

1801 CGCTACGTGTGCGAGTGCGCTCGGGGCTACGGCGGCTCAACTGCCAGTTCTGCTCCCC 1860

1861 GAGCCACCTCAGGCGCCGGTCATCGTTGACTTCACCGAGAACTACACAGAGGGCCAGAAC 1920

1921 AGCCAGTTTCCCTGGATCGCAGTGTCGCGCGGGATTATTCTGGTCCTCATGCTGCTGCTG 1980

FIG. 1A (cont'd)

2401 TACCAGTCGGTGTACGTCATATCAGAAGAGAAAGATGAGTGCCATCATAGCAACTGAGGTG 2460

2461 TAAACAGACGTGACGTGGCAAAGCTTATCGATACCGTCATCAAGCTT 2508

FIG. 1A (cont'd)

[illegible]

1 GAATTCGGCAGGAGGTTTTTTTTTTTTTCCCTCTTTCTTTCTTTCTTTTCCATCCGAAAG 69

70 AGCTGTCAGCCGCCGCCGGGCTGCACCTAAAGGCGTCGGTAGGGGGATAACAGTCAGAGACCTCCTGA 138

139 AAGCAGGAGACGGGACGGTACCCCTCCGGCTCTGCGGGCGGCTGCGGCCCTCCGTCTTTCCCCCTC 207

208 CCCGAGAGACACTCTTCTTTCCCCCACGAAGACACAGGGGCAGGAACCGAGCGCTGCCCCCTCCGCC 276

277 ATGGGAGGCCGCTTCTCTGCTGACGCTCGCCCTCTCTCGGCGCTGCTGTGCGGCTCCAGGTTGACGGC 345

346 TCCGGGGTGTTTCGAGCTGAAGCTGCAGGAGTTGTCAACAAGAAGGGGCTGCTCAGCAACCGCAACTGC 414

415 TGCCGGGGGGGGCGGCCCGGAGGCGCGGGCAGCAGTGGGACTGCAAGACCTTCTTCCGCGTCTGC 483

FIG. 1B

484 CTGAAGCACTACCAGGCCAGCGTCTCCCCGAGCCGCCCTGCACCTACGGCAGCGCCAT

553 CTCGGCGCCAACCTCCTTCAGCGTCCCGACGGCGCGGGCGGGCGCCGACCCCGCCTTCA

622 CGCTTCCCCTTCGGCTTCACCTGGCCCGGCACCTTCTCGCTCATCATCGAGGCTCTCC

691 CCCGACGACCTCACCACAGAAAACCCCGAGCGCCTCATCAGCCGCCTGGCCACCCAGA

760 GTGGGCGAGGAGTGGTCCCAGGACCTGCACAGCAGCGGCCGCACTGACCTCAAGTAC

FIG. 1B (cont'd)

GTGTGATGAGCACTACTACGGGAAGGCTGCTCTGTCTTCTGCGGCCCCGTGACGACCGCTTCGGT 897

898(CACTTCACCTGTGGAGAGCGTGGCGAGAAGGTCTGCAACCCAGGCTGGAAAGGCCAGTACTGCACTGAG 966

967 CCGATTTCCTTGCCTGGGTGTGACGAGCAGCACGGCTTCTGCGACAAACCTGGGGAATGCAAGTGCAGA 1035

1036 GTGGGTGGCAGGGGCGGTACTGTGACGAGTGCATCCGATACCCAGGCTGCCTGCACGGTACCTGTCAG 1104

1105 CAGCCATGGCAGTGCAACTGCCAGGAAGGCTGGGGCGGCCTTTCTGCAACCAGGACCTGAACTACTGC 1173

1174 ACTCACCACAAGCCATGCAAGAATGGTGCCACATGCACCAACACCGGTCAGGGGAGCTACACTTGTCT 1242

1243 TGCCGACCTGGGTACACAGGCTCCAGCTGCGAGATTGAAATCAACGAATGTGATGCCAACCCTTGCAAG 1311

FIG. 1B (cont'd)

1312 AATGGTGGAAGCTGCACGGATCTCGAGAACAGCTATTCTGTACCTGCCCCCAGGCTTCTATGGTAAA 1380

1381 AACTGTGAGCTGAGTGCAATGACTTGTGCTGATGGACCGTGCTTCAATGGAGGGCGATGCACTGACAAC 1449

1450 CCTGATCGTGGATACAGCTGCCGCTGCCCACTGGGTTATTCTGGGTTCAACTGTGAAAAGAAAATCGAT 1518

1519 TACTGCAGTTCCAGCCCTTGTGCTAATGGAGCCCAGTGCGTTGACCTGGGGAACCTACATATGCCAG 1587

1588 TGCCAGGCTGGCTTCACTGGCAGGCACTGTGACGACAACGTGGACGATTGCGCCTCCTTCCCCTGCGTC 1656

1657 AATGGAGGGACCTGTCAGGATGGGGTCAACGACTACTCCTGCACCTGCCCCCGGGATACAACGGGAAG 1725

FIG. 1B (cont'd)

1726 AACTGCAGCACGCCGGTGAGCAGATGCGAGCACAACCCCTGCCACAATGGGGCCACCTGCCACGAGAGA 1794

1795 AGCAACCGCTACGTGTGCGAGTGCCTCGGGGCTACGGCGGCCTCAACTGCCAGTTCTGCTCCCCGAG 1863

1864 CCACCTCAGGGGCGGTCATCGTTGACTTCACGAGAAGTACACAGAGGGCCAGAACAGCCAGTTTCCC 1932

1933 TGGATCGCAGTGTGCGCCGGGATTATTCTGGTCCTCATGCTGCTGCTGGGTTGCGCCGCCATCGTCGTC 2001

2002 TCGGTCAGGCTGAAGGTGCAGAAGAGGCCACCAACAGCCGAGGCTGCAGGAGTGAAACGGAGACCATG 2070

2071 AACAACTGGCGAACTGCCAGCGCGAGAAGGACATCTCCATCAGCGTCATCGGTGCCACTCAGATTAAA 2139

2140 AACACAAATAAGAAAGTAGACTTTTCACAGCGATAACTCCGATAAAAACGGCTACAAAGTTAGATACCCA 2208

2209 TCAGTGGATTACAATTGTTGGTGCATGAACTCAAGAATGAGGACTCTGTGAAAGAGGAGCATGGCAAATGC 2277

2278 GAAGCCAAGTGTGAAACGTATGATTCAGAGGCAGAAGAGAAAAGCGCAGTACAGCTAAAAAGTAGTGAC 2346

2347 ACTTCTGAAAGAAAACGGCCAGATTCAGTATATTCCACTTCAAAGGACACAAAGTACCAGTCGGTGTAC 2415

2416 GTCATATCAGAAGAGAAAGATGAGTGCATCATAGCAACTGAGGTTAGTATCCCACTGGCAGTCGGACA 2484

2485 AGTCTTGGTGTGTGATTCCCATCTAGCGCAGGTGAGGGCGGCCAAACCATTCCTACCTGCTGCCACAGTC 2553

2554 ATCTGTACCCAATGAAAACCTGGCCACCTTCAGTCTGTGGCACTGCAGACGTTGAAAAAAGTTGTTGTGG 2622

FIG. 1B (cont'd)

2623 ATTAPCATAAGCTCCAGTGGGGGTTACAGGGACAGCAATTTTTCAGGCAAGGGTATAACTGTAGTGCA 2691

2692 GTGTAGCTTACTAACCTACTGACTCATTCTTTCGTGTCTTCCTGCAGAGCCTGTTTTTGCTTGGCA 2760

2761 TTGAGGTGAAGTCCTGACCCTCTGCATCCTCATAGTCCTCTGCTTCTTTTTATTAAACCTCTTCTGGTC 2829

2830 TCTGCTTGCTTTTCTCTCAACAGGTGTAAACAGACGTGACGTGGCAAGCTT 2883

FIG. 1B (cont'd)

1 MGRFLLTLA LLSALLCRCQ VDGSGVFELK LOEFVNKKGL LSNRNCCRGG GPGGAGQQQC
61 DCKTFFRVCL KHYQASVSPE PPCTYGSALT PVLGANSFSV PDGAGGADPA FSNPIRFPFG
121 FTWPGTFSLI IEALHTDSPD DLTENPERL ISRLATQRHL AVGEESQDL HSSGRTDLKY
181 SYRFVCDEHY YGEGCSVFCR PRDDRFGHFT CGERGEKVCN PGWKGQYCTE PICLPGCDEQ
241 HGFCDKPGEC KCRVGWQGRY CDECIRYPGC LHGTCQQPWQ CNCQEGWGL FCNQDLNYCT
301 HHKPCKNAT CTNTGQGSYT CSCRPGYTGS SCEIEINECD ANPCKNGGSC TDLENSYSCT
361 CPPGFYGNK ELSAMTCADG PCFNGGRCTD NPDGGYSCRC PLGYSGFNCE KKIDYCSSSP
421 CANGAQCVDL GNSYICQQA GFTGRHCDDN VDDCASFCV NGGTCQDGVN DYSCTCPPGY
481 NGKNCSTPVS RCEHNPCHNG ATCHERSNRY VCECARGYGG LNCQFLLPEP PQGPVIVDFT
541 EKYTEGQNSQ FPWIAVCAGI ILVLMLLGC AAIIVCVRLK VQKRHHQPEA CRSETETMNN
601 LANCQREKDI SISVIGATQI KNTNKKVDFH SDNSDKNGYK VRYPVSVDYNL VHELKNEDSV
661 KEEHGKCEAK CETYDSEAE KSAVQLKSSD TSEKRPDSV YSTSKDTKYQ SVYVISEEKD
721 ECIIATEV

FIG. 2

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C-Delta-1	1	MGGRFLTLA-LLSALLCRCOVDGSGVPELKLOEFVNKKGLLSNRNCCRGGGPGGAGQQQC	60
X-Delta-1	1	MGQORMLTLL-VLSAVL--COISCSGLFELRLOEFVNKKGLLGNMNCCRPQSL--ASLQRC	56
Delta	1	--MHWIKCLLTAFICFTVIVQVHSSGSFELRLKYFSNDHGRDNEGRCCSGESDGA TGKCLG	59
C-Delta-1	61	DCKTFFRVCLKHYQASVSPEPPCTYGSAITPVLGANSFVSVPDAGGADPAFSPNPIRFPFGF	121
X-Delta-1	57	ECKTFFRICLKHYQSNVSPEPPCTYGGAVTPVLGTNSPVVPESSNADPTFSPNPIRFPFGF	116
Delta	60	SKTRFRRLCLKHYQATIDTTSQCTYGDVITPILGENSVNLTDAQRFQNKGETNPIQFPFSE	120
C-Delta-1	122	TWPGTFSLIIEALHTDSPDDLTTENPERLISRLATQRHLAVGREWSQDLHSSGRTDLKYSY	182
X-Delta-1	117	TWPGTFSLIIEAIIHADSAADDLNTENPERLISRLATQRHLTVGEQWSQDLHSSDRTELKYSY	177
Delta	121	SWPGTFSLIVEAWH-DTNNSGNARTNKLIIQRLLVQVLEVSSEWKTNKSESQYTSLEYDF	180
C-Delta-1	183	RFVCDHEYHYGEGCSVFCRPRDDRFGHFTCGERGEKVCNPGWKQYCTEPICLPGCDEQHGF	243
X-Delta-1	178	RFVCDHEYHYGEGCSDYCRPRDDAFGHFSCGERGEKLCNPGWKGLYCTEPICLPGCDEHHGY	238
Delta	181	RVTCDLNYYGSGCAKFCRPRDDSFHSTCSSTGEIICLTGWQGDYCHIPKCAKGC--HGH	239
		DSL	
C-Delta-1	244	CDKPGECKCRVGVWQGRYCDECIRYPGCLHGTCQQPWCNCQEGWGGGLFCNQDLNYCTHKKP	304
X-Delta-1	239	CDKPGECKCRVGVWQGRYCDECIRYPGCLHGTCQQPWCNCQEGWGGGLFCNQDLNYCTHKKP	299
Delta	240	CDKPNQCVCQLGWKGAALCNECVLEPNCTIHGTCNKPWTICINEGWGGGLFCNQDLNYCTHHRP	300
		EGF1	EGF2
C-Delta-1	305	CKNGATCTNTGQGSYTCSCRPGYTGSCEIEINECDA--NPCKNGGSCTD--LENSYSCT	360
X-Delta-1	300	CENGAATCTNTGQGSYTCSCRPGYTGSNCEIEVNECDA--NPCKNGGSCSD--LENSYSCS	355
Delta	301	CKNGGTCTPNTGEGLYTCKCAPGYSGDDCENEIYSCDAVDNPCQNGGTCIDEPHTKTGYKCH	361
		EGF3	EGF4
C-Delta-1	361	CPPGFYGNCELSAMTCADGPCFNG-----GRCTDNPDDGGYSRCPLGYSGFNCEKKIDYC	416
X-Delta-1	356	CPPGFYGNCELSAMTCADGPCFNG-----GRCA DNPDDGGYICFCPGVYSGFNCEKKIDYC	411
Delta	362	CRNGWSGKMCEKVLTCSDKPCHQGICRNVVRPGLSGKGQGYQCECPIGYSGFNCIDLQLDNC	422
		EGF5	
C-Delta-1	417	SSSPCANGAQCVDLGNSYICOCQAGFTGRHCDDNVDDCASFPFCVNGGTCODGVNDYSCTCP	477
X-Delta-1	412	SSNPCANGARCEDLGNSYICOCQEGFSGRNCDDNLD DCTSFPCQNGGTCODGI NDYSCTCP	472
Delta	423	SPNPCIINGGSCQPSGK--CICP SGFSGTRCETNIDDC LGHQCENGGTCIDMVNQYRCQCV	480
		EGF6	EGF7
C-Delta-1	478	PGYNGRNCSTPVSRCEHNPNCHNGATCHERSNRYVCECARGYGGLLNCQFLLPEPPQGP----	534
X-Delta-1	473	PGYIGKNCSTPITKCEHNPNCHNGATCHERNRYVCCARGYGGNNCQFLLPE-----	524
Delta	481	PGFHHGTHCSKVDLCLIRPCANGGTCLNLNNDYQCTCRAGFTGKDCSV DIDECS SGPC HNG	541
		EGF8	
C-Delta-1	535	-----VIVDFTE--KYTEGQNSQPPW--IAVCAGIILVL	564
X-Delta-1	525	-----EKPVVVDLTE--KYTEGQSGQPPW--IAVCAGIIVLV	557
Delta	542	GTCMNRVNSPECVCANGFRGKQCD EESYDSVTFDAHQYGATTQARADGLANAQVVLIAVFS	602
		EGF9	
C-Delta-1	565	MLLLGCAAIVVCVRLKVKQRHHQPEACRS ETE TMNNLANCQREKD--ISISVIGATQIKNT	623
X-Delta-1	558	MLLLGCAAIVVCVVRVVKRRRHQPEACRGE SK TMNNLANCQREKD--ISVSFIGTTOIKNT	616
Delta	603	VAMP LVAVIAACVVFCHMKRRKRAQEKDNAEARKQNEQNAVATMHHNGSAVGVALASASMG	663
		TM	
C-Delta-1	624	NKKVDFHSD--NSDKNGYKVRYPSPVDYNLVHELKNEDSVKEEHGKCEAKCETYDSEAEKSA	683
X-Delta-1	617	NKKIDFLSESNNKNGYKPRYPSPVDYNLVHELKNEDSPKEERSKCEAKCSSNDSDSE DVNS	677
Delta	664	GKTGSNSGLTFDGGNPNIIKNTWDKSVN-NICASAAAAAAAAAAAADECLHYGGYVASVADN	723
C-Delta-1	684	-----VQLKSSDTSERK-----RPDSVYSTSKDTKYQSVYVISEBKDECIATEV	728
X-Delta-1	678	-----VHSK-RDSSERK-----RPDSAYSTSKDTKYQSVYVISDEKDECIATEV	721
Delta	724	NNANSDFCVAPLQRAKSQKQLNTDPTLMHRGSPAGTSAKGASGGGPGAAEGKRI SVLGEGS	784
Delta	785	YCSQRWPSLAAAGVAGACSSQLMAAASAAAGTDGTAQQQRSVVCGTPHM	832

FIG. 3

C-Delta-1	184	V-CDEHYIYGE	G-CSVFCRPR	DDRFGEFTCG	ERGEKVCNPG	WKGQYC	228
Delta	182	VTCDLNYYGS	G-CAKFCRPR	DDSFGEHSTCS	ETGEIICLTG	WQGDYC	226
Serrate	235	VQCAVTYYNT	TFCTTFCRPR	DDQFGHYACG	SEGQKLCNLG	WQGVNC	279
C-Serrate-1		VTCAEHYYGF	G-CNKFCRPR	DDFFTHHTCD	QNGNKTCLG	WTGPEC	
apx-1	130	NLCSSNYHGK	R-CNRYCIAN	-AKLHWE-CS	THGVRRCSAG	WSGEDC	172
lag-2	120	VTCAARNYFGN	R-CENFCDAH	LAKAARKRCD	AMGRLRCDIG	WMGPHC	166

FIG. 4

Country	Year	Population (millions)	Urban population (millions)	Urban population (%)	Population density (per sq km)	Urban population density (per sq km)	Population growth rate (%)	Urban population growth rate (%)	Population growth rate (%)	Urban population growth rate (%)	Population growth rate (%)	Urban population growth rate (%)
Algeria	1980	12.5	4.5	36	100	100	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	1985	13.5	5.5	41	110	110	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	1990	14.5	6.5	45	120	120	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	1995	15.5	7.5	48	130	130	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2000	16.5	8.5	52	140	140	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2005	17.5	9.5	54	150	150	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2010	18.5	10.5	57	160	160	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2015	19.5	11.5	59	170	170	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2020	20.5	12.5	61	180	180	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2025	21.5	13.5	63	190	190	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2030	22.5	14.5	64	200	200	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2035	23.5	15.5	66	210	210	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2040	24.5	16.5	67	220	220	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2045	25.5	17.5	69	230	230	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2050	26.5	18.5	70	240	240	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2055	27.5	19.5	71	250	250	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2060	28.5	20.5	72	260	260	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2065	29.5	21.5	73	270	270	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2070	30.5	22.5	74	280	280	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2075	31.5	23.5	75	290	290	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2080	32.5	24.5	76	300	300	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2085	33.5	25.5	76	310	310	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2090	34.5	26.5	77	320	320	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2095	35.5	27.5	77	330	330	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2100	36.5	28.5	78	340	340	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2105	37.5	29.5	79	350	350	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2110	38.5	30.5	79	360	360	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2115	39.5	31.5	80	370	370	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2120	40.5	32.5	80	380	380	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2125	41.5	33.5	81	390	390	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2130	42.5	34.5	81	400	400	1.5	1.5	1.5	1.5	1.5	1.5
Algeria	2135	43.5	35.5									

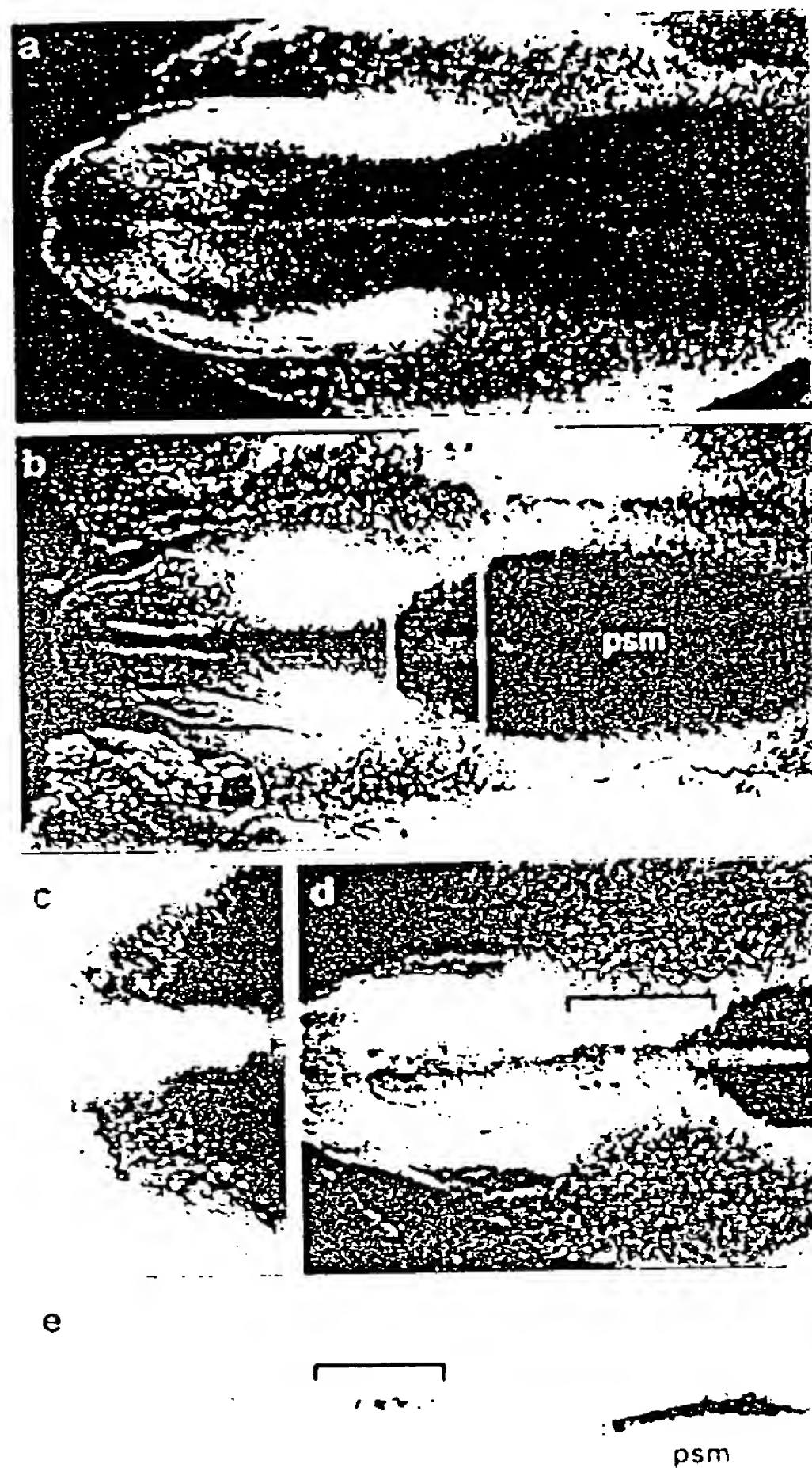


FIG. 5

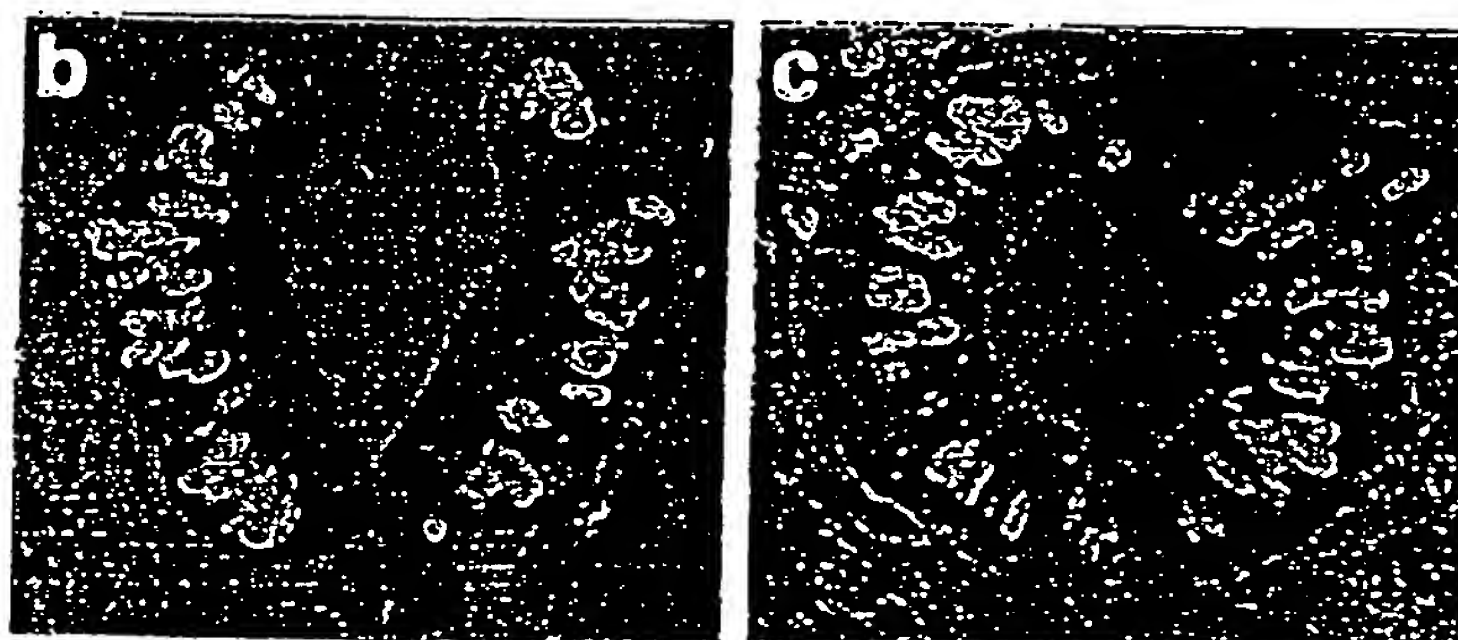


FIG. 6B

FIG. 6C

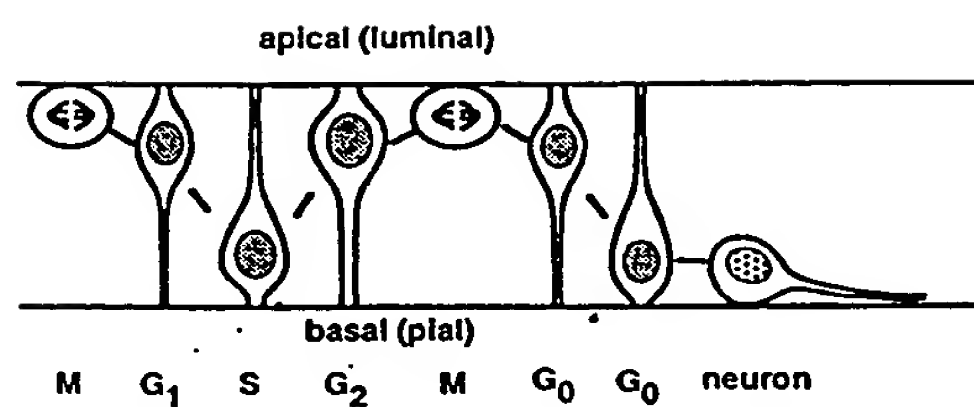


FIG. 6A

CTGCAGGAAT TCSMYCGCAT GCTCCCGGCC GCCATGGGCC GTCGGAGCGC GCTAGCCCTT 60
 GCCGTGGTCT CTGCCCTGCT GTGCCAGGTC TGGAGCTCCG GCGTATTTGA GCTGAAGCTG 120
 CAGGAGTTCG TCAACAAGAA GGGGCTGCTG GGAACCGCA ACTGCTGCCG CGGGGGCTCT 180
 GGCCCGCCTT GCGCCTGCAG GACCTTCTTT CCGGTATGCC TCAAGCACTA CCAGGCCAGC 240
 GTGTCACCGG AGCCACCCTG CACCTACGGC AGTGCCGTCA CGCCAGTGCT GGGTGTGAC 300
 TCCTTCAGCC TGCCTGATGG CGCAGGCATC GACCCCGCCT TCAGCAACCC CATCCGATTC 360
 CCCTTCGGCT TCACCTGGCC AGGTACCTTC TCTCTGATCA TTGAAGCCCT CCATACAGAC 420
 TCTCCCGATG ACCTCGCAAC AGAAAACCCA GAAAGACTCA TCAGCCGCCT GACCACACAG 480
 AGGCACCTCA CTGTGGGAGA AGAATGGTCT CAGGACCTTC ACAGTAGCGG CCGCACAGAC 540
 CTCCGGTACT CTTACCGGTT TGTGTGTGAC GAGCACTACT ACGGAGAAGG TTGCTCTGTG 600
 TTCTGCCGAC CTCGGGATGA CGCCTTTGGC CACTTCACCT GCGGGGACAG AGGGGAGAAG 660
 ATGTGCGACC CTGGCTGGAA AGGCCAGTAC TGCCTGACC CAATCTGTCT GCCAGGGTGT 720
 GATGACCAAC ATGGATACTG TGACAAACCA GGGGAGTGCA AGTGACAGAGT TGGCTGGCAG 780
 GGCCGCTACT GCGATGAGTG CATCCGATAC CCAGGTTGTC TCCATGGCAC CTGCCAGCAA 840
 CCCTGGCAGT GTAAC TGCCA GGAAGGCTGG GGGGGCCTTT TCTGCAACCA AGACCTGAAC 900
 TACTGTACTC ACCATAAGCC GTGCAGGAAT GGAGCCACCT GCACCAACAC GGGCCAGGGG 960
 AGCTACACAT GTTCCTGCCG ACCTGGGTAT ACAGGTGCCA ACTGTGAGCT GGAAGTAGAT 1020
 GAGTGTGCTC CTAGCCCCTG CAAGAACGGA GCGAGCTGCA CGGACCTTGA GGACAGCTTC 1080
 TCTTGACCTT GCCCTCCCGG CTTCTATGGC AAGGTCTGTG AGCTGAGCGC CATGACCTGT 1140
 GCAGATGGCC CTTGCTTCAA TGGAGGACGA TGTTTCAGATA ACCCTGACGG AGGCTACACC 1200
 TGCCATTGCC CCTTGGGCTT CTCTGGCTTC AACTGTGAGA AGAAGATGGA TCTCTGCGGC 1260
 TCTTCCCCTT GTTCTAACGG TGCCAAGTGT GTGGACCTCG GCAACTCTTA CCTGTGCCGG 1320
 TGCCAGGCTG GCTTCTCCGG GAGGTACTGC GAGGACAATG TGGATGACTG TGCCCTCCTCC 1380
 CCGTGTGCAA ATGGGGGCAC CTGCCGGGAC AGTGTGAACG ACTTCTCCTG TACCTGCCCA 1440
 CCTGGCTACA CGGGCAAGAA CTGCAGCGCC CCTGTGAGCA GGTGTGAGCA TGCACCCTGC 1500
 CATAATGGGG CCACCTGCCA CCAGAGGGGC CAGCGCTACA TGTGTGAGTG CGCCAGGGC 1560
 TATGGCGGCC CCAACTGCCA GTTTCTGCTC CCTGAGCCAC CACCAGGGCC CATGGTGGTG 1620
 GACCTCAGTG AGAGGCATAT GGAGAGCCAG GCGGGGCCCT TCCCCTGGGT GGCCGTGTGT 1680
 GCCGGGGTGG TGCTTGTCCT CCTGCTGCTG CTGGGCTGTG CTGCTGTGGT GGTCTGCGTC 1740
 CGGCTGAAGC TACAGAAACA CCAGCCTCCA CCTGAACCCT GTGGGGGAGA GACAGAAACC 1800
 ATGAACAACC TAGCCAATTG CCAGCGCGAG AAGGACGTTT CTGTTAGCAT CATTGGGGCT 1860
 ACCCAGATCA AGAACACCAA CAAGAAGGCG GACTTTCACG GGGACCATGG AGCCGAGAAG 1920
 AGCAGCTTTA AGGTCCGATA CCCCCTGTG GACTATAACC TCGTTCGAGA CCTCAAGGGA 1980
 GATGAAGCCA CGGTCAGGGA TACACACAGC AAACGTGACA CCAAGTGCCA GTCACAGAGC 2040
 TCTGCAGGAG AAGAGAAGAT CGCCCCAACA CTTAGGGGTG GGGAGATTCC TGACAGAAAA 2100
 AGGCCAGAGT CTGTCTACTC TACTTCAAAG GACACCAAGT ACCAGTCGGT GTATGTTCTG 2160
 TCTGCAGAAA AGGATGAGTG TGTTATAGCG ACTGAGGTGT AAGATGGAAG CGATGTGGCA 2220
 AAATTCCCCT TTCTCTTAAA TAAAATTCCA AGGATATAGC CCCGATGAAT GCTGCTGAGA 2280
 GAGGAAGGGA GAGGAAACCC AGGGACTGCT GCTGAGAACC AGGTTTCAGGC GAACGTGGTT 2340
 CTCTCAGAGT TAGCAGAGGC GCCCGACACT GCCAGCCTAG GCTTTGGCTG CCGCTGGACT 2400
 GCCTGCTGGT TGTTCCTTGC GCACTATGGA CAGTTGCTTT GAAGAGTATA TATTTAAATG 2460
 GACGAGTGAC TTGATTCATA TAGGAAGCAC GCACTGCCCA CACGTCTATC TTGGATTACT 2520
 ATGAGCCAGT CTTTCCTTGA ACTAGAAACA CAACTGCCTT TATGTCTCTT TTTGATACTG 2580
 AGATGTGTTT TTTTTTTTTC CTAGACGGGA AAAAGAAAAC GTGTGTTATT TTTTTTGGGA 2640
 TTTGTAAAAA TATTTTTCAT GATTATGGGA GAGCTCCCAA CCGTTGGAG GT 2692

FIG. 7

MGRRSALALA	VVSALLCQVW	SSGVFELKLQ	EFVNKKGLLG	NRNCCRGGSG	50
PPCACRTFFR	VCLKHYQASV	SPEPPCTYGS	AVTPVLGVDS	FSLPDGAGID	100
PAFSNPIRFP	FGFTWPGTFS	LIIEALHTDS	PDDLATENPE	RLISRLTTQR	150
HLTVGEEWSQ	DLHSSGRSDL	RYSYRFVCDE	HYYGEGCSVF	CRPRDDAFGH	200
FTCGDRGEKM	CDPGWKQYQ	TDPICLPGCD	DQHGCDKPG	ECKCRVGWQG	250
RYCDECIRYP	GCLHGTCQQP	WQCNCQEGWG	GLFCNQDLNY	CTHHKPCRNG	300
ATCTNTGQGS	YTCSCRPGYT	GANCELEVDE	CAPSPCKNGA	SCTDLED\$FS	350
CTCPPGFYBK	VCELSAMTCA	DGPCFNGGRC	SDNPDGGYTC	HCPLGFSGFN	400
CEKKMDLCGS	SPCSNGAKCV	DLGNSYLCRC	QAGFSGRYCE	DNVDDCASSP	450
CANGGTCRDS	VNDFSCTCPP	GYTGKNCSAP	VSRCEHAPCH	NGATCHQRGQ	500
RYMCECAQGY	GGPNCQFLLP	EPPPGPMVVD	LSEHMHESQG	GPFPWVAVCA	550
GVVLVLLLLL	GCAAVVVCVR	LKLQKHQPPP	EPCGGETETM	NNLANCQREK	600
DVSVSIIGAT	QIKNTNKKAD	FHGDHGAES	SFKVRYPTVD	YNLVRDLKGD	650
EATVRDTHSK	RDTKCQSQSS	AGEEKIAPTL	RGGEIPDRKR	PESVYSTSKD	700
TKYQSVYVLS	AEKDECVIAT	EV			722

FIG. 8

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Chick DELTA	YGERFLITAYLLSAILCRQ	YDCSGVFEELKLOEFVNRKGLRL	NRNCGRGG	50													
Mouse Delta.pep	YGRSALMIAVVSAILCQ--	YNSGVFEELKLOEFVNRKGLRL	NRNCGRGG	48													
Consensus	YGRN...IAV...SAILC...	Y...SGVFEELKLOEFVNRKGLRL	NRNCGRGG	50													
Chick DELTA	GPGGAGQQQZKQK	YDFERVCLKHVOASVSPEPPCTVGSAT	TPVLGANSSEV	100													
Mouse Delta.pep	--SGP---PCTNCR	YDFERVCLKHVOASVSPEPPCTVGSAT	TPVLGANSSEV	93													
Consensus	...S...PCTNCR	YDFERVCLKHVOASVSPEPPCTVGSAT	TPVLGANSSEV	100													
Chick DELTA	PDGAGSADPAESNDRPPEFG	ETWPGTFESLIT	TEAHTDSDP	150													
Mouse Delta.pep	PDGAG-IDPAESNDRPPEFG	ETWPGTFESLIT	TEAHTDSED	142													
Consensus	PDGAG...IDPAESNDRPPEFG	ETWPGTFESLIT	TEAHTDSED	150													
Chick DELTA	TSRI	TORHL	YGEWSODL	HSSGRDLE	200												
Mouse Delta.pep	TSRI	TORHL	YGEWSODL	HSSGRDLE	192												
Consensus	TSRI	TORHL	YGEWSODL	HSSGRDLE	200												
Chick DELTA	PRDD	EGHET	CG	RGEK	250												
Mouse Delta.pep	PRDD	EGHET	CG	RGEK	242												
Consensus	PRDD	EGHET	CG	RGEK	250												
Chick DELTA	KCRVWOGRY	CDECIRYPGC	LHGTCCOPWQ	CNCOEGWGG	FCNODLNYCT	300											
Mouse Delta.pep	KCRVWOGRY	CDECIRYPGC	LHGTCCOPWQ	CNCOEGWGG	FCNODLNYCT	292											
Consensus	KCRVWOGRY	CDECIRYPGC	LHGTCCOPWQ	CNCOEGWGG	FCNODLNYCT	300											
Chick DELTA	HHKPC	NGAT	CTNTGCGSYT	CSCRPGYTG	SC	EE	NECD	ANPCKNG	SC	350							
Mouse Delta.pep	HHKPC	NGAT	CTNTGCGSYT	CSCRPGYTG	SC	EE	NECD	ANPCKNG	SC	342							
Consensus	HHKPC	NGAT	CTNTGCGSYT	CSCRPGYTG	SC	EE	NECD	ANPCKNG	SC	350							
Chick DELTA	TDLE	S	SCT	CPPGFYK	C	ELSAMTCADG	PCFNGGRC	D	NPDGGY	SC	400						
Mouse Delta.pep	TDLE	S	SCT	CPPGFYK	C	ELSAMTCADG	PCFNGGRC	D	NPDGGY	SC	392						
Consensus	TDLE	S	SCT	CPPGFYK	C	ELSAMTCADG	PCFNGGRC	D	NPDGGY	SC	400						
Chick DELTA	PLG	SGFNCE	KK	D	C	SSP	C	NGA	CVDL	GNSY	C	COA	GF	GR	C	DN	450
Mouse Delta.pep	PLG	SGFNCE	KK	D	C	SSP	C	NGA	CVDL	GNSY	C	COA	GF	GR	C	DN	442
Consensus	PLG	SGFNCE	KK	D	C	SSP	C	NGA	CVDL	GNSY	C	COA	GF	GR	C	DN	450
Chick DELTA	VDDCAS	PCV	NGGTC	DS	VN	D	SCTCPPGY	NGKNCS	PVS	RCEH	PCHNG	500					
Mouse Delta.pep	VDDCAS	PCV	NGGTC	DS	VN	D	SCTCPPGY	NGKNCS	PVS	RCEH	PCHNG	492					
Consensus	VDDCAS	PCV	NGGTC	DS	VN	D	SCTCPPGY	NGKNCS	PVS	RCEH	PCHNG	500					
Chick DELTA	ATCH	ERSNRY	VECAR	GYGG	NCOFFLLPEP	P	GEV	VDFT	EKYTE	ONSQ	550						
Mouse Delta.pep	ATCH	ERSNRY	VECAR	GYGG	NCOFFLLPEP	P	GEV	VDFT	EKYTE	ONSQ	542						
Consensus	ATCH	ERSNRY	VECAR	GYGG	NCOFFLLPEP	P	GEV	VDFT	EKYTE	ONSQ	550						
Chick DELTA	EPW	AVCAG	I	VVL	LLGC	AA	VVCVRLK	OKRHQPEA	RSSETETMNN	600							
Mouse Delta.pep	EPW	AVCAG	I	VVL	LLGC	AA	VVCVRLK	OKRHQPEA	RSSETETMNN	592							
Consensus	EPW	AVCAG	I	VVL	LLGC	AA	VVCVRLK	OKRHQPEA	RSSETETMNN	600							
Chick DELTA	LANCOREKDI	S	S	IGATQI	KNTNKK	DFH	SDN	SDK	GY	KVRYE	VDYN	649					
Mouse Delta.pep	LANCOREKDI	S	S	IGATQI	KNTNKK	DFH	SDN	SDK	GY	KVRYE	VDYN	642					
Consensus	LANCOREKDI	S	S	IGATQI	KNTNKK	DFH	SDN	SDK	GY	KVRYE	VDYN	650					
Chick DELTA	LV	HEKNE	D	SVKEB	KCE	AKCETYDSEA	EEKSA	VOLKS	S	TSERKRPD	698						
Mouse Delta.pep	LV	HEKNE	D	SVKEB	KCE	AKCETYDSEA	EEKSA	VOLKS	S	TSERKRPD	692						
Consensus	LV	HEKNE	D	SVKEB	KCE	AKCETYDSEA	EEKSA	VOLKS	S	TSERKRPD	700						
Chick DELTA	SVYSTSKDTK	YOSVYV	S	E	KDEC	LATEV	728										
Mouse Delta.pep	SVYSTSKDTK	YOSVYV	S	E	KDEC	LATEV	722										
Consensus	SVYSTSKDTK	YOSVYV	S	E	KDEC	LATEV	730										

FIG. 9

[illegible]

FIG. 10

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90	90	90	90	90
91	91	91	91	91
92	92	92	92	92
93	93	93	93	93
94	94	94	94	94
95	95	95	95	95
96	96	96	96	96
97	97	97	97	97
98	98	98	98	98
99	99	99	99	99
100	100	100	100	100

FIG. 10 (cont'd)

[illegible]

FIG. 11

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(SHEET 25 OF 37)

10	20	30	40	50	60
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
CATTGGGTAC	GGGCCCCCT	CGAGGTCGAC	GGTATCGATA	AGCTTGATAT	CGAATTCCGG
70	80	90	100	110	120
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
CTTCACCTGG	CCGGGCACCT	TCTCTCTGAT	TATTGAAGCT	CTCCACACAG	ATTCTCCTGA
130	140	150	160	170	180
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
TGACCTCGCA	ACAGAAAACC	CAGAAAGACT	CATCAGCCGC	CTGGCCACCC	AGAGGCACCT
190	200	210	220	230	240
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
GACGGTGGGC	CAGGAGTGGT	CCCAGUACCT	GCACAGCAGC	GGCCGCACGG	ACCTCAAGTA
250	260	270	280	290	300
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
CTCCTACCGC	TTCGTGTGTC	ACGAACACTA	CTACGGAGAG	GGCTGCTCCG	TTTCTOCCG
310	320	330	340	350	360
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
TCCCCGGGAC	GATGCCTTCG	GCCACTTCAC	CTGTGGGGAG	CGTGGGGAGA	AAGTGTGCAA
370	380	390	400	410	420
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
CCCTGGCTGG	AAAGGGCCCT	ACTGCACAGA	GCCGATCTGC	CTGCCTGGAT	GTGATGAGCA
430	440	450	460	470	480
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
GCATGGATTT	TGTGACAAAC	CACCGGAATC	CAAGTGCAGA	G'TGGGCTGGC	AGGGCCGGTA
490	500	510	520	530	540
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
CTGTGACGAG	TGTATCCGCT	ATCCAGGCTG	TCTCCATGGC	ACCTGCCAGC	AGCCCTGGCA
550	560	570	580	590	600
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
GTGCAACTGC	CAGGAAGGNT	GGGGGCCCCT	TTTCTGCAAC	CAGGACCTGA	ACTACTGCAC
610	620	630	640	650	660
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
ACACCATAAG	CCCTGCAAGA	ATGGAGCCAC	CTGCAACAAA	CACGGGCCAG	GGGGAGCTAC
670	680	690	700	710	720
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
ACTTGGTCTT	TGGCCGGNCT	GGGGTACANA	GGGTGCCACC	TGCGAAGCTT	GGGGATTGGA
730	740	750	760	770	780
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
CGAGTTGTTG	ACCCAGCCC	TTGGTAAGAA	CGGAGGGAGC	T'TGACGGA'IC	TTCGGAGAAC
790	800	810	820	830	840
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
AGCTACTCCT	GTACCTGCCC	ACCCGGCTTC	TACGUCAAAA	TCT'IG'IGAA'I'I	GAGTGCCATG
850	860	870	880	890	900
* * * *	* * * *	* * * *	* * * *	* * * *	* * * *
ACCTGTGCGG	ACGGCCCTTG	CTT'TAACGGG	GGTCCGTGCT	CAGACAGCCC	CGATGGAGGG

FIG. 12A

910	920	930	940	950	960
* * *	* * *	* * *	* * *	* * *	* * *
TACAGCTGCC	GCIGCCCCGT	GGGCTACTCC	GGCTTCAACT	GTGAGAAGAA	AATTGACTAC
970	980	990	1000	1010	1020
* * *	* * *	* * *	* * *	* * *	* * *
TGCAGCTCTT	CACCCGTGTC	TAATGGTCCC	AAGTGTGTGG	ACCTCGGTGA	TGCCTACCTG
1030	1040	1050	1060	1070	1080
* * *	* * *	* * *	* * *	* * *	* * *
TGCCGCTGCC	AGGCCGGCTT	CTCGGGGAGG	CACTGTGACG	ACAACGTGGA	CGACTGCGCC
1090	1100	1110	1120	1130	1140
* * *	* * *	* * *	* * *	* * *	* * *
TCCTCCCCGT	GCGCCAACGG	GGGCACCTGC	CGGGATGGCG	TGAACGACTT	CTCCTGCACC
1150	1160	1170	1180	1190	1200
* * *	* * *	* * *	* * *	* * *	* * *
TGCCCCGCTG	GCTACACGGG	CAGGAACCTG	AGTGCCCCCG	CCAGCACOTG	CGAGCACGCA
1210	1220	1230	1240	1250	1260
* * *	* * *	* * *	* * *	* * *	* * *
CCCTGCCACA	ATGGGGCCAC	CTGCCACGAG	AGGGGCCACC	GCTATNTGTG	CGAGTGTGTC
1270	1280	1290	1300	1310	1320
* * *	* * *	* * *	* * *	* * *	* * *
CGAAGCTACG	GGGGTCCCAA	CTUCCANTTC	CTGCTCCCCC	AAACTGCCCC	CCCGGCCCCA
1330	1340	1350	1360	1370	1380
* * *	* * *	* * *	* * *	* * *	* * *
CGGTGGTGGA	AAC'TCCCCTA	AAAAAACCTA	AAAGGGCCGG	GGGGGGCCCA	TCCCC'TTGGT
1390	1400	1410	1420	1430	1440
* * *	* * *	* * *	* * *	* * *	* * *
GGACGTGTGC	GCCGGGGTCA	TCC'TGTCTT	CATGCTGCTC	CTGGGCTGTG	CCGCTGTGGT
1450	1460	1470	1480	1490	1500
* * *	* * *	* * *	* * *	* * *	* * *
GGTCTGCGTC	CGGCTGAGGC	TGCAGAAGCA	CCGGCCCCCA	CCCGACCCCT	GNCGGGGGGA
1510	1520	1530	1540	1550	1560
* * *	* * *	* * *	* * *	* * *	* * *
GACGGAGACC	ATGAACAACC	TGGNCAACTG	CCAGCGTIGAG	AAGGACATCT	CAGTCAGCAT
1570	1580	1590	1600	1610	1620
* * *	* * *	* * *	* * *	* * *	* * *
CATCGGGGNC	ACGCAGATCA	AGAACACCAA	CAAGAAGGCG	GACTTCCACG	GGGACCACAG
1630	1640	1650	1660	1670	1680
* * *	* * *	* * *	* * *	* * *	* * *
NGCCGACAAG	AATGGCTTCA	AGGCCCCGTA	CCAGNGGTG	GACTATAACC	TCGTGCAGGA
1690	1700	1710	1720	1730	1740
* * *	* * *	* * *	* * *	* * *	* * *
CC'TCAAGGGT	GACGACACCG	CCGTCAGCOA	CGCGCACAGC	AAGCGTGACA	CCAAGTGNCA
1750	1760	1770	1780	1790	1800
* * *	* * *	* * *	* * *	* * *	* * *
GCCCCAGGGC	TCCTCAGGGG	AGGACAAAGG	GACCCCCGAC	CCACACTCAG	GGGGTGGAGG
1810	1820	1830	1840	1850	1860
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FIG. 12A (cont'd)

AAGCATCTTG AAAGAAAAG GCCGGACTTC GGGCTTG'TC AACTTTCAA AGACAANCAA
 1870 1880 1890 1900 1910 1920
 * * * * * * * * * *
 NGTACAAGTC GGTGT'NCGTC ATTTCCGNAG GAGGAAGGNT GACT'GCGTCA TAGGAANTTG
 1930 1940 1950 1960 1970 1980
 * * * * * * * * * *
 AGGTNG'IAAA NTGGNAGTTG ANN'I'GGAAA GNNNTCCCCG GATTCGNTT TCAAAGTTTT
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FIG. 12A (cont'd)

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CATGGGTAC GGGCCCCCT CGAGGTCGAC GGTATCGATA AGCTTGATAT CGAATTCGGG
H W V R A P L E V D G I D K L D I E F R> 20
I G Y G P P S R S T V S I S L I S N S [G] 20
L G T G P P R G R R Y R * A * Y R I P> 19

70 80 90 100 110 120

CTTCACCTGG CCGGGCACCT TCTCTCTGAT TATTGAAGCT CTCCACACAG ATTCTCCTGA
L H L A G H L L S D Y * S S P H R F S *> 40
F T W P G T F S L I I E A L H T D S P D> 40
A S P G R A P S L * L L K I S T Q I L L> 39

130 140 150 160 170 180

TGACCTCGCA ACAGAAAACC CAGAAAGACT CATCAGCCGC CTGGCCACCC AGAGGCACCT
* P R N R K P R K T H Q P P G H P E A P> 60
D L A T E N P E R L I S R L A T Q R H L> 60
M T S Q Q K T Q K D S S A A W P P R - G T> 59

190 200 210 220 230 240

GACGGTGGC GAGGAG'GGT' CCCAGGACCT GCACAGCAGC GGCCGCACGG ACCTCAAGTA
D G G R G V V P G P A Q Q R P H G P Q V> 80
T V G E E W S Q D L H S S G R T D L K Y> 80
* R W A R S G P R T C T A A A A R T S S> 79

250 260 270 280 290 300

CTCCTACCGC TTCGTGTGTG ACGAACACTA CTACGGAGAG GGCTGCTCCG TTTTCTGCCG
L L P L R V * R T L L R R G L L R F L P> 100
S Y R F V C D E H Y Y G E G C S V F C R> 100
T P T A S C V T N T T T E R A A P F S A> 99

310 320 330 340 350 360

TCCCCGGGAC GATGCCTTCG GCCACTTCAC CTGTGGGGAG CGTGGGGAGA AAGTGTGCAA
S P G R C L R P L H L W G A W G E S V Q> 120
P R D D A F G H F T C G E R G E K V C N> 120
V P G T M P S A T S P V C S V G R K C A> 119

370 380 390 400 410 420

CCCTGGCTGG AAAGGGCCCT ACTGCACAGA GCCGATCTGC CTGCCTGGAT GTGATGAGCA
P W L E R A L L H R A D L P A W M * * A> 140
P G W K G P Y C T E P L C L P G C D E Q> 140
T I A G K G P T A Q S R S A C L D V M S> 139

430 440 450 460 470 480

GCATGGATTT TGTGACAAAC CAGCCCAATG CAAGTGCAGA GTGGGCTGGC AGGGCCGGTA
A W I L * Q T R G M Q V Q S G L A G P V> 160
H G F C D K P G E C K C R V G W Q G R Y> 160
S M D F V T N Q G N A S A E W A G R A G> 159

490 500 510 520 530 540

CTGTGACGAG TGTATCCGCT ATCCAGGCTG TCTCCATGGC ACCTGCCAGC AGCCCTGGCA
L * R V Y P L S R L S P W H L P A A L A> 180

FIG. 12B

C D E C I R Y P G C L H G T C Q Q P W O> 180
 T V T S V S A I Q A V S M A P A S S P O> 179
 550 560 570 580 590 600
 * * * * *
 GIGCAACTGC CAGGAAGGNT GGGGGGGCCT TTTCTGCAAC CAGGACCTGA ACTACTGCAC
 V Q L P G R X G G P F L O P G P E L L H> 200
 C N C Q E G W G G L F C N Q D L N Y C T> 200
 S A T A R K X G G A F S A T R T * T T A> 199
 610 620 630 640 650 660
 * * * * *
 ACACCATAAG CCCTGCAAGA ATCGAGCCAC CTGCAACAAA CACGGGCCAG GGGGAGCTAC
 T P * A L Q E W S H L Q Q T R A R G S Y> 220
 H H K P C K N G A T C N K H G P G C A T> 220
 H T I S P A R M E P P A T N T G Q G E L> 219
 670 680 690 700 710 720
 * * * * *
 ACTTGGTCTT TGGCCGGNCT GGGGTACANA GGGTGCCACC TCGAAGCTT GGGGATTGGA
 T W S L A G L G Y X G C H L R S L G I G> 240
 L G L W P X W G T X G A T C E A W G L D> 240
 H L V F G R X C V X R V P P A K L G D W> 239
 730 740 750 760 770 780
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 CGAGTTGTG ACCCCAGCCC TTGTAAGAA CGGAGGGAGC TTGACGGATC TTCGGAGAAC
 R V V D P S P W * E R R E L D G S S E N> 260
 E L L T P A L G K N G G S L T D L R R T> 260
 T S C * P Q P L V R T E O A * R I F G E> 259
 790 800 810 820 830 840
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 AGCTACTCCT GTACCTGCCC ACCCGGCTTC TACGGCAAAA TCIGTGAATT GAGTGCCATG
 S Y S C T C P P G F Y G K I C E L S A M> 280
 A T P V P A H P A S T A K S V N * V P *> 280
 Q L L L Y L P T R L L R Q N L * I E C H> 279
 850 860 870 880 890 900
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 ACCTGTGCGG ACGGCCCTTG CTTTAAACGGG GGTGCGTGCT CACACAGCCC CGATGGAGGG
 T C A D G P C F N G G R C S D S P D G G> 300
 P V R T A L A L T G V G A Q T A P M E G> 300
 D L C G R P L L * R G S V I R Q P R W R> 299
 910 920 930 940 950 960
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 TACAGCTGCC GCTGCCCCGT GGGCTACTCC GCCTTCAACT GTGAGAAGAA AATTGACTAC
 Y S C R C P V G Y S G F N C E K K I D Y> 320
 T A A A A P W A T P A S T V R R K L T T> 320
 V Q L P L P R G L L R L Q L * E E N * I> 319
 970 980 990 1000 1010 1020
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 TGCAGCTCTT CACCCTGTC TAATGGTGG AAGTGTGTGG ACCTGGGTGA TGCCTACCTG
 C S S S P C S N G A K C V D L G D A Y L> 340
 A A L H P V L M V P S V W T S V M P T C> 340
 I Q L F T L F * W C Q V C G P R * C L P> 339
 1030 1040 1050 1060 1070 1080
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 TGCCGCTGCC AGGCCGGCTT CTCGCCGAGG CACTGTGACG ACAACGTGGA CGACTGCGCC

FIG. 12B (cont'd)

C R C Q A G F S G R H C D D N V D D C A> 360
 A A A R P A S R G G T V T T T W T T A P> 360
 V P L P G R L L G F A L * R Q R G R L R> 359
 1090 1100 1110 1120 1130 1140
 * * * * *
 TCCCTCCCGT GCGCCAACGG GGGCACCTGC CCGGATGGCG TGAACGACTT CTCCTGCACC 380
 S S P C A N G G T C R D G V N D F S C T> 380
 P P R A P T G A P A G M A * T T S P A P> 380
 L L P V R Q R G H L P G W R E R I L L H> 379
 1150 1160 1170 1180 1190 1200
 * * * * *
 TGCCCGCCTG GCTACACGGG CAGGAAGTGC AGTGGCCCCG CCAGCAGGTC CGAGCAGGCA 400
 C P P G Y T G R N C S A P A S R C E H A> 400
 A R L A T R A G T A V P P P A G A S T H> 400
 L P A W L H G Q E L Q C P R Q Q V R A R> 399
 1210 1220 1230 1240 1250 1260
 * * * * *
 CCCTGCCACA ATGGGUCCAC CTGCCACGAG AGGGGCCACC GCTATATGTG CGAGTGTGCC 420
 P C H N G A T C H F R G H R Y X C E C A> 420
 P A T M G P P A T R G A T A J C A S V P> 420
 T L P Q W G H L P R E G P P L F V R V C> 419
 1270 1280 1290 1300 1310 1320
 * * * * *
 CGAAGCTACG GGGGTCCCAA CTGCCATTC CTGCTCCCGG AAAGTGGCCC CCGGGCCCCA 440
 R S Y G G P N C X F L L P E T A P P A P> 440
 F A T G V P T A X S C S P K L P P R P H> 440
 P K L R G S Q L P X P A P R N C P P G P> 439
 1330 1340 1350 1360 1370 1380
 * * * * *
 CGGTGGTGGG AACTCCCCTA AAAAAACCTA AAAGGGCCGG GGGGGGCCCA TCCCCTTGGT 460
 R W W K L P * K N L K G P G G A H P L G> 460
 G G G N S P K K T * K G R G G P I P L V> 460
 T V V F T P L K K P K R A G G G P S P W> 459
 1390 1400 1410 1420 1430 1440
 * * * * *
 GGACGTGTGC GCCGGGGTCA TCCCTGTCCT CATGCTGCTG CTGGGCTGTC CCGCTGTGGT 480
 G R V R R G H P C P H A A A G L C R C G> 480
 D V C A G V I L V L M L L L G C A A V V> 480
 W T C A P G S S L S S C C C W A V P L W> 479
 1450 1460 1470 1480 1490 1500
 * * * * *
 GGTCTGCGTC CCGCTGAGGC TGCAGAAGCA CCGGCCXXCA GCCGACCCCT GNCGGGGGGA 500
 G L R P A E A A E A P A P S R P L X G G> 500
 V C V R L R L Q K H R P P A D P X R G E> 500
 W S A S G * G C R S T G P Q P T P X G G> 499
 1510 1520 1530 1540 1550 1560
 * * * * *
 GACGGAGACC ATGAACAACC TGGNCAACIG CCAGCGTGAG AAGGACATCT CAGTCAGCAT 520
 D C D H E Q P C Q L P A * E G H L S Q H> 520
 T E T M N N L X N C Q R E K D I S V S I> 520
 R R R P * T T W X T A S V R R T S Q S A> 519
 1570 1580 1590 1600 1610 1620
 * * * * *

FIG. 12B (cont'd)

CATCGGGGNC ACGCAGATCA AGAACACCAA CAAGAAGGCG GACTTCCACG GGGACCACAG
 H R G H A D Q E H Q Q E G G L P R G P Q> 540
 I G X T O I K N T N K K A D F H G D H X> 540
 S S G X R R S R T P T R R R T S T Q T T> 539
 1630 1640 1650 1660 1670 1680
 * * * * *
 NGCCGACAAG AATGGCTTCA AGGCCCGCTA CCCAGNCGTG GACTATAACC TCGTGCAGGA
 X R Q E W L Q G P L P X G G L * P R A G> 560
 A D K N G F K A R Y P X V D Y N L V Q D> 560
 X P T R M A S R P A T Q X W T I T S C R> 559
 1690 1700 1710 1720 1730 1740
 * * * * *
 CCTCAAGGCT GACGACACCG CCGTCAGGGA CGCGCACAGC AAGCGTGACA CCAAGTGNCA
 P Q G * R H R R Q G R A Q Q A * H Q V X> 580
 L K G D D T A V R D A H S K R D T K X Q> 580
 T S R V T T P P S G T R T A S V T P S X> 579
 1750 1760 1770 1780 1790 1800
 * * * * *
 GCCCCAGGGC TCCTCAGGGG AGGAGAAGGG GACCCCGAC CCACACTCAG GGGGTGGAGG
 A P G L L R G G H Q D P R P T L R G W R> 600
 P O G S S G E E K G T P D P H S G G G G> 600
 S P R A P Q G R R R C P P T H T Q G V R> 599
 1810 1820 1830 1840 1850 1860
 * * * * *
 AAGCATCTTG AAAGAAAAAG GCCGGACTTC GGGCTTGTTT AACTTTCAAA AGACAANCAA
 K H L E R K R P D F G L V Q L S K D X Q> 620
 S I L K E K G R T S G L F N F Q K T X X> 620
 E A S * K K K A G L R A C S T F K R Q X> 619
 1870 1880 1890 1900 1910 1920
 * * * * *
 NGTACAAGTC GGTGTNCGTC ATTTCCGNAG GAGGAAGGNT GACTGCGTCA TAGGAANTIG
 X T S R C X S F P X E E G * L R H R X L> 640
 V Q V G V R H F R R R K X D C V T G X *> 640
 X Y K S V X V I S X G G R X T A S * E X> 639
 1930 1940 1950 1960 1970 1980
 * * * * *
 AGGTNGTAAA NTGGNAGTTG ANNTKGA AAA GNNNTCCCCO GATTCCGNTT TCAAAGTTTT
 R X * X G S * X W K X X P G F R F Q S F> 660
 G X K X X V X X G K X S P D S X F K V F> 660
 E V V X W X L X L E X X P R I P X S K F> 659

T

FIG. 12B (cont'd)

Mouse Delta vs Partial Human Delta

Mouse Delta DNA	GTCCAGCGGT ACCATGGGCC GTCCGAGCGC GCTACCCCTT GCGTGGTCT	50
Human Delta	-----	
Consensus	GTCCAGCGGT ACCATGGGCC GTCCGAGCGC GCTACCCCTT GCGTGGTCT	50
Mouse Delta DNA	CTGCCCTGCT GTGCCAGGTC TGGAGCTCCG GCGTATTGA GCTGAAGCTG	100
Human Delta	-----	
Consensus	CTGCCCTGCT GTGCCAGGTC TGGAGCTCCG GCGTATTGA GCTGAAGCTG	100
Mouse Delta DNA	CAGGAGTTCG TCAACAAGAA GGGGCTGCTG GGAACCCCA ACTGCTGCCG	150
Human Delta	-----	
Consensus	CAGGAGTTCG TCAACAAGAA GGGGCTGCTG GGAACCCCA ACTGCTGCCG	150
Mouse Delta DNA	CGGGGGCTCT GGGCCGCTT GCGCTGCAG GACCTTCTT CCGTATGCC	200
Human Delta	-----	
Consensus	CGGGGGCTCT GGGCCGCTT GCGCTGCAG GACCTTCTT CCGTATGCC	200
Mouse Delta DNA	TCAAGCACTA CCAGGCCAGC GTGTACCGG AGCCACCCTG CACCTACGGC	250
Human Delta	-----	
Consensus	TCAAGCACTA CCAGGCCAGC GTGTACCGG AGCCACCCTG CACCTACGGC	250
Mouse Delta DNA	AGTGCTGTCA CGCCAGTGCT GGGTGTGAC TCCTTCAGCC TGCCTGATG	300
Human Delta	-----	5
Consensus	AGTGCTGTCA CGCCAGTGCT GGGTGTGAC TCCTTCAGCC TGCCTGATG	300
Mouse Delta DNA	CGGAGGCAIC GACCTC--G GGTTCAGAA CCGA--TCC GATTC--CCC	343
Human Delta	GGTACGGGC CCGCTCGAGG ICCACGGTAT CGATAGCTT GATPTCGAAT	55
Consensus	SGAGGCRYC SMCCTCGAGG TCKTCGMAW CSMYAGYYY GATPTCGMY	350
Mouse Delta DNA	TCGGCTTCA CCTGGCCGG TACCTTCTCT CTGATATTG AAGCTCTCCA	393
Human Delta	TCGGCTTCA CCTGGCCGG TACCTTCTCT CTGATATTG AAGCTCTCCA	105
Consensus	TCGGCTTCA CCTGGCCGG TACCTTCTCT CTGATATTG AAGCTCTCCA	400
Mouse Delta DNA	TACAGATCTT CCTGATGACC TCGCAACAGA AAACCCAGAA AGACTCATCA	443
Human Delta	CACAGATCTT CCTGATGACC TCGCAACAGA AAACCCAGAA AGACTCATCA	155
Consensus	TACAGATCTT CCTGATGACC TCGCAACAGA AAACCCAGAA AGACTCATCA	450
Mouse Delta DNA	GCGGCTGAC CACACAGAGG CACCTACCG TGGGAGAGA TTGGTCCAG	493
Human Delta	GCGGCTGAC CACACAGAGG CACCTACCG TGGGAGAGA TTGGTCCAG	205
Consensus	GCGGCTGAC CACACAGAGG CACCTACCG TGGGAGAGA TTGGTCCAG	500
Mouse Delta DNA	GACCTCACA GAGCGGCCG CACGACCTC CGTACTCTT ACCGTTTGT	543
Human Delta	GACCTCACA GAGCGGCCG CACGACCTC AGTACTCTT ACCGTTTGT	255
Consensus	GACCTCACA GAGCGGCCG CACGACCTC CGTACTCTT ACCGTTTGT	550
Mouse Delta DNA	GTGTGACGAG CACTACTACG GAGAGGGTGT CTCGTCTTC TGCCGACCTC	593
Human Delta	GTGTGACGAG CACTACTACG GAGAGGGTGT CTCGTCTTC TGCCGACCTC	305
Consensus	GTGTGACGAG CACTACTACG GAGAGGGTGT CTCGTCTTC TGCCGACCTC	600
Mouse Delta DNA	GGGATGATGC CTTGGCCAC TTCACCTGG GGGACAGGG GGAGAGATG	643
Human Delta	GGGATGATGC CTTGGCCAC TTCACCTGG GGGACAGGG GGAGAAAGTG	355
Consensus	GGGATGATGC CTTGGCCAC TTCACCTGG GGGASMGAG GGAGAAARTG	650

FIG. 13

Mouse Delta vs Partial Human Delta

Mouse Delta DNA	TGCCACCCCTG GCTGGAAAGG	CCAGTACTGC ACAGACCCA TCTGCTGCC	693
Human Delta	TGCCACCCCTG CCTGGAAAGG	CCAGTACTGC ACAGACCCA TCTGCTGCC	405
Consensus	TGCCACCCCTG GCTGGAAAGG	CCAGTACTGC ACAGACCCA TCTGCTGCC	700
Mouse Delta DNA	AGGTTGTGAT GACCAACATG GATACTGTGA CAAACCAGGG GATTGCAAGT		743
Human Delta	TGGTTGTGAT GACCAACATG GATTGTGTGA CAAACCAGGG GAATTGCAAGT		455
Consensus	AGGTTGTGAT GACCAACATG GATTGTGTGA CAAACCAGGG GAATTGCAAGT		750
Mouse Delta DNA	GCAGAGTGG CTGGCAGGGC CGTACTGCG ATGAGTGTAT CCGTATCCA		793
Human Delta	GCAGAGTGG CTGGCAGGGC CGTACTGTG ATGAGTGTAT CCGTATCCA		505
Consensus	GCAGAGTGG CTGGCAGGGC CGTACTGTG ATGAGTGTAT CCGTATCCA		600
Mouse Delta DNA	GGTTGTCTCC ATGGCACCTG CCAGCACCC TGGCAGTGA ACTGCCAGGA		843
Human Delta	GGTTGTCTCC ATGGCACCTG CCAGCACCC TGGCAGTGA ACTGCCAGGA		555
Consensus	GGTTGTCTCC ATGGCACCTG CCAGCACCC TGGCAGTGA ACTGCCAGGA		850
Mouse Delta DNA	AGGTTGGGGG GGCCTTTTCT GCAACCA GA CCTGAACTAC TGTACTCACC		893
Human Delta	AGGTTGGGGG GGCCTTTTCT GCAACCA GA CCTGAACTAC TGTACTCACC		605
Consensus	AGGTTGGGGG GGCCTTTTCT GCAACCA GA CCTGAACTAC TGTACTCACC		900
Mouse Delta DNA	ATAAGCCCTG CAGGAATGGA GCCACCTGCA CCAACACGG GCCAGGGG A		941
Human Delta	ATAAGCCCTG CAGGAATGGA GCCACCTGCA CCAACACGG GCCAGGGG A		655
Consensus	ATAAGCCCTG CAGGAATGGA GCCACCTGCA CCAACACGG GCCAGGGG A		950
Mouse Delta DNA	GCTACACATG ATCTT-GCC GACCTGGGT ATATA GGTG CCAACTGTG-		986
Human Delta	GCTACACATG ATCTTTGGCC GACCTGGGT ATATA GGTG CCAACTGTG		705
Consensus	GCTACACATG ATCTTTGGCC GACCTGGGT ATATA GGTG CCAACTGTG		1000
Mouse Delta DNA	AGCT--GGGA GTAGATGAG TG-TGCTCTT AGCCCTTGC AAGAACGGAG		1031
Human Delta	AGCTTGGGA GTAGATGAG TGTGTGACCTT AGCCCTTGGT AAGAACGGAG		755
Consensus	AGCTTGGGA GTAGATGAG TGTGTGACCTT AGCCCTTGGT AAGAACGGAG		1050
Mouse Delta DNA	CGAGCTTCAC GGAICTT--G AGACAGCTT CTCTTGACC TGCCCTCCCG		1079
Human Delta	CGAGCTTCAC GGAICTTGG AGACAGCTT CTCTTGACC TGCCCTCCCG		605
Consensus	CGAGCTTCAC GGAICTTGG AGACAGCTT CTCTTGACC TGCCCTCCCG		1100
Mouse Delta DNA	GCTTCTAAGG CAAGTCTGT GAGCTGACCG CCATGACCTG TGCAGATGGC		1129
Human Delta	GCTTCTAAGG CAAGTCTGT GAATGTAGTG CCATGACCTG TGCAGATGGC		855
Consensus	GCTTCTAAGG CAAGTCTGT GAGCTGACCG CCATGACCTG TGCAGATGGC		1150
Mouse Delta DNA	CCTTGCTTAA AAGGAGGCG ATGTCAGAT ACCCTGACG GAGGTACAC		1179
Human Delta	CCTTGCTTAA AAGGAGGCG ATGTCAGAT ACCCTGACG GAGGTACAG		905
Consensus	CCTTGCTTAA AAGGAGGCG ATGTCAGAT ACCCTGACG GAGGTACAG		1200
Mouse Delta DNA	CTGCCATTCG CCGTTGGGCT CTCTGGCTT CAACTGTGAG AAGAAATGG		1229
Human Delta	CTGCCATTCG CCGTTGGGCT CTCTGGCTT CAACTGTGAG AAGAAATGG		955
Consensus	CTGCCATTCG CCGTTGGGCT CTCTGGCTT CAACTGTGAG AAGAAATGG		1250
Mouse Delta DNA	ATCTCTGCG CTCTTCCTT TGTCTAAGG GTGCCAAGTG TGTGGACCTC		1279
Human Delta	ATCTCTGCG CTCTTCCTT TGTCTAAGG GTGCCAAGTG TGTGGACCTC		1005
Consensus	ATCTCTGCG CTCTTCCTT TGTCTAAGG GTGCCAAGTG TGTGGACCTC		1300

FIG. 13 (cont'd)

Mouse Delta vs Partial Human Delta

Mouse Delta DNA	GGCAACTCCT ACCTGTGCCG TCCCAAGCT GGCTTCTCG GGAGGACTG	1329
Human Delta	GGTGATCCCT ACCTGTGCCG TGGCAGGCC GGCTTCTCG GGAGGACTG	1055
Consensus	GGTATTCCT ACCTGTGCCG TGGCAGGCC GGCTTCTCG GGAGGACTG	1350
Mouse Delta DNA	CGAGGACAA GTGGATGACT GGGCTCCTC CCCGTGAGCA AATGGGGCA	1379
Human Delta	TGAGGACAA GTGGATGACT GGGCTCCTC CCCGTGAGCC AATGGGGCA	1105
Consensus	TGAGGACAA GTGGATGACT GGGCTCCTC CCCGTGAGCA AATGGGGCA	1400
Mouse Delta DNA	CCTGCCGGGA CAGGTGAAC GACTTCTCT GACCTGCCC CCTGGCTAC	1429
Human Delta	CCTGCCGGGA TGGGTGAAC GACTTCTCT GACCTGCCC CCTGGCTAC	1155
Consensus	CCTGCCGGGA TGGGTGAAC GACTTCTCT GACCTGCCC CCTGGCTAC	1450
Mouse Delta DNA	ACGGGCAGA ACTGCAG GC CCCAGCAGC AGGTGAGC AAGCACCTG	1479
Human Delta	ACGGGCAGA ACTGCAG GC CCCAGCAGC AGGTGAGC AAGCACCTG	1205
Consensus	ACGGGCAGA ACTGCAG GC CCCAGCAGC AGGTGAGC AAGCACCTG	1500
Mouse Delta DNA	CCATAATGGG GCCACCTGCC AAGAGAGGG CCAACGCTAC ATGTGTGAGT	1529
Human Delta	CCATAATGGG GCCACCTGCC AAGAGAGGG CCAACGCTAC ATGTGTGAGT	1255
Consensus	CCATAATGGG GCCACCTGCC AAGAGAGGG CCAACGCTAC ATGTGTGAGT	1550
Mouse Delta DNA	GGGCCAGGG CTATGGGGG CCCAACTGCC AATTCTGCT CCGTGAAGC	1578
Human Delta	GAGGCCAAG CTATGGGGG CCCAACTGCC AATTCTGCT CCGTGAAGC	1305
Consensus	GAGGCCAAG CTATGGGGG CCCAACTGCC AATTCTGCT CCGTGAAGC	1600
Mouse Delta DNA	-ACCCAGGG CCCCATGGTG GTGGATCTC AGTGATAGGC ATATGCAGA	1625
Human Delta	GCCCCAGGG CCCCATGGTG GTGGATCTC CCTAATAAA ACCTAATAGG	1355
Consensus	GACCCAGGG CCCCATGGTG GTGGATCTC MSYATARRM AATTAATAGG	1650
Mouse Delta DNA	GGCAGGGGG GGGCTTCCCC TTCTCTGCG TGTGAGCCGG GGTGCTCTT	1675
Human Delta	GGCAGGGGG GGGCTTCCCC TTCTCTGCG TGTGAGCCGG GGTGCTCTT	1405
Consensus	GGCAGGGGG GGGCTTCCCC TTCTCTGCG TGTGAGCCGG GGTGCTCTT	1700
Mouse Delta DNA	GTCCTCTGC TGCTGCTGGG CTGTGCGCT GTGGTGGTCT GCGTCCGGCT	1725
Human Delta	GTCCTCTGC TGCTGCTGGG CTGTGCGCT GTGGTGGTCT GCGTCCGGCT	1455
Consensus	GTCCTCTGC TGCTGCTGGG CTGTGCGCT GTGGTGGTCT GCGTCCGGCT	1750
Mouse Delta DNA	GAGGCTCAG AAACACCGGC CACCACTGA ACCCTGTGGG GGAGAGACG	1775
Human Delta	GAGGCTCAG AAACACCGGC CACCACTGA ACCCTGTGGG GGAGAGACG	1505
Consensus	GAGGCTCAG AAACACCGGC CACCACTGA ACCCTGTGGG GGAGAGACG	1800
Mouse Delta DNA	AAACCATGAA CAACCTGAC AATGCCAGC GGAGAGAAG CATTCTGTT	1825
Human Delta	AAACCATGAA CAACCTGAC AATGCCAGC GGAGAGAAG CATTCTGTT	1555
Consensus	AAACCATGAA CAACCTGAC AATGCCAGC GGAGAGAAG CATTCTGTT	1850
Mouse Delta DNA	AGCATCATG GGGTACCA GATCAAGAAC ACCAACAAGA AGGCGGACTT	1875
Human Delta	AGCATCATG GGGTACCA GATCAAGAAC ACCAACAAGA AGGCGGACTT	1605
Consensus	AGCATCATG GGGTACCA GATCAAGAAC ACCAACAAGA AGGCGGACTT	1900
Mouse Delta DNA	TCACGCGGAC CATGGAGCA AGAAGACAG CTTAAGGTC CGTACCCCA	1925
Human Delta	CCACCCGAC CATGGAGCA AGAAGATAG CTTAAGGTC CGTACCCCA	1655
Consensus	TCACGCGGAC CATGGAGCA AGAAGARYG CTTAAGGTC CGTACCCMR	1950

FIG. 13 (cont'd)

Mouse Delta vs Partial Human Delta

Mouse Delta DNA	CTGTGGACTA TAACCTCGTT	CCAGACCTCA AGGGAGAGA	AGCCACCGTC	1975
Human Delta	NGGTGGACTA TAACCTCGTG	CAGGACCTCA AGGGAGAGA	TAACCTCGTC	1705
Consensus	NKGTGGACTA TAACCTCGTK	CRGACCTCA AGGGAGAGA	TRCCACCGTC	2000
Mouse Delta DNA	AGGGATACAC ACAGCAACCG	TGACACCAAG TGACAGTAC	AGAGCTCTGC	2025
Human Delta	AGGGACCCCG ACAGCAACCG	TCACACCAAG TGCAGCTCC	AGAGCTCTTC	1755
Consensus	AGGGATTCAC ACAGCAACCG	TGACACCAAG TGCAGCTCC	AGAGCTCTTC	2050
Mouse Delta DNA	AGGAGAGGAG AA--GATCG	CC--CCACA CTGA--GGGGT	GG--GG--AGAT	2067
Human Delta	ACGGAGAGGAG AAGGGGACCC	CCGACCTACA CTGA--GGGGT	GGAGGAGAGTA	1805
Consensus	AGGAGAGGAG AAGGGGACCS	CCGACCTACA CTGA--GGGGT	GGAGGAGAGW	2100
Mouse Delta DNA	TCGTGAGAGA AAAAGGCCCG	ACTCT--GTC TACTCTAC	T TCAAAAGGAC	2113
Human Delta	TCGTGAGAGA AAAAGGCCCG	ACTTCGGGCT TGTTCACIT	TCAAAAGACA	1855
Consensus	TCGTGAGAGA AAAAGGCCCG	ACTTYGGGY TRVTCACIT	TCAAAAGACA	2150
Mouse Delta DNA	-ACCAAGTAC CAGTCGGTGT	ATGTTCTCTC TGAGAA--A	AGGATGAGTG	2160
Human Delta	ANCAAGTAC CAGTCGGTGT	NGTCAATTC CGAGGAGGA	AGGATGAGTG	1905
Consensus	ANCAAGTAC CAGTCGGTGT	NYGTYMTKTC MNAAGAGGA	AGGATGAGTG	2200
Mouse Delta DNA	TGTATATA--GC GACTGAGCT-	GTAAGATGGA AGCGATCTGS	CAAAATTCCC	2208
Human Delta	CGTCATAGGA ANTTGAGGTN	GTAANTTGGN AG--T-TG--	--ANNTT---	1945
Consensus	YGTATATAGM RNYTGAGCTN	GTAARNITGN AGCGATCTGS	CAANNTTCCC	2250
Mouse Delta DNA	ATTCTCTCTCA AATAAAATTC	CAAGGATATA GCCCGATGA	ATGCTCTCTGA	2258
Human Delta	-----GGA AAGNNN- TC	CCGGAT-- --TCCGTT--	----TTC---	1972
Consensus	ATTCTCTCKSA AAKNNNATTC	CMCGATATA GCYCCGNTGA	ATGCTCTCTGA	2300
Mouse Delta DNA	GAGAGGAAGG GAGAGCAAA:	CCAGGGACTG CTGCTGAGAA	CCAGGTTTCAG	2308
Human Delta	-- -- -- -- --AAA--	-- -- -- -- --GTTT--	-- -- -- -- --	1981
Consensus	GAGAGGAAGG GAGAGCAAA:	CCAGGGACTG MKVITGAGAA	CCAGGTTTCAG	2350
Mouse Delta DNA	GCGAAGCTGG TTCTCTCAGA	GTAGCAGAG GCGCCCGACA	CTGCCAGCCT	2358
Human Delta	-- -- -- -- --	-- -- -- -- --	-- -- -- -- --	1981
Consensus	GCGAAGCTGG TTCTCTCAGA	GTTAGCAGAG GCGCCCGACA	CTGCCAGCCT	2400
Mouse Delta DNA	AGGCTTTGGC TGCCCTTGA	CTGCTGCTG GITGTTCCCA	TTGCACTATG	2408
Human Delta	-- -- -- -- --	-- -- -- -- --	-- -- -- -- --	1981
Consensus	AGGCTTTGGC TGCCCTTGA	CTGCTGCTG GITGTTCCCA	TTGCACTATG	2450
Mouse Delta DNA	GACAGTTGCT TTGAAGAGTA	TATATTTAAA TGGACGAGTG	ACTTGATTCA	2458
Human Delta	-- -- -- -- --	-- -- -- -- --	-- -- -- -- --	1981
Consensus	GACAGTTGCT TTGAAGAGTA	TATATTTAAA TGGACGAGTG	ACTTGATTCA	2500
Mouse Delta DNA	TATAGGAAGC ACGCACTGCC	CACACGTCTA TCTTGATTA	CTATGAGCCA	2508
Human Delta	-- -- -- -- --	-- -- -- -- --	-- -- -- -- --	1981
Consensus	TATAGGAAGC ACGCACTGCC	CACACGTCTA TCTTGATTA	CTATGAGCCA	2550
Mouse Delta DNA	GTCCTTCCTT GAACTAGAAA	CACAACTGCC TTTATTGTCC	TTTTTGATAC	2558
Human Delta	-- -- -- -- --	-- -- -- -- --	-- -- -- -- --	1981
Consensus	GTCCTTCCTT GAACTAGAAA	CACAACTGCC TTTATTGTCC	TTTTTGATAC	2600

FIG. 13 (cont'd)

Mouse Delta vs Partial Human Delta

Mouse Delta DNA	TGAGATGTGT	TTTTTTT	TTTT	CCTAGACGGG	AAAAAGAAAA	CCTGTGTTAT	2608
Human Delta	-----	-----	-----	-----	-----	-----	1981
Consensus	TGAGATGTGT	TTTTTTTTT		CCTAGACGGG	AAAAAGAAAA	CGTGTGTTAT	2650
Mouse Delta DNA	TTTTTTGGGA	TTTGTA AAAA	TATTTTTCAT	GATATCTGTA	AAGCTTGACT		2658
Human Delta	-----	-----	-----	-----	-----	-----	1981
Consensus	TTTTTTGGGA	TTTGTA AAAA	TATTTTTCAT	GATATCTGTA	AAGCTTGACT		2700
Mouse Delta DNA	ATTTTGTGAC	GTTTCATTTT	TTATAATTTA	AATTTTGGTA	AATATGTACA		2708
Human Delta	-----	-----	-----	-----	-----	-----	1981
Consensus	ATTTTGTGAC	GTTTCATTTT	TTATAATTTA	AATTTTGGTA	AATATGTACA		2750
Mouse Delta DNA	AAGGCACTTC	GGGTCTATGT	GACTATATTT	TTTGTATAT	AAATGTATTT		2758
Human Delta	-----	-----	-----	-----	-----	-----	1981
Consensus	AAGGCACTTC	GGGTCTATGT	GACTATATTT	TTTGTATAT	AAATGTATTT		2800
Mouse Delta DNA	ATGGAATATT	GTGCAAAATGT	TATTTGAGTT	TTTACTGTT	TTGTTAATGA		2808
Human Delta	-----	-----	-----	-----	-----	-----	1981
Consensus	ATGGAATATT	GTGCAAAATGT	TATTTGAGTT	TTTACTGTT	TTGTTAATGA		2850
Mouse Delta DNA	AGAAATTCAT	TTTAAAAATA	TTTTTCCAAA	ATAAATATAA	TGAACTACA		2857
Human Delta	-----	-----	-----	-----	-----	-----	1981
Consensus	AGAAATTCAT	TTTAAAAATA	TTTTTCCAAA	ATAAATATAA	TGAACTACA		2899

FIG. 13 (cont'd)

G F T W P G T F S L I I E A L H T D S P D> 21
 D L A T E N P E R L I S R L A T Q R H L> 41
 T V G E E W S Q D L H S S G R T D L K Y> 61
 S Y R F V C D E H Y Y G E G C S V F C R> 81
 P R D D A F G H F T C G E R G E K V C N> 101
 P G W K G P Y C T E P I C L P G C D E Q> 121
 H G F C D K P G E C K C R V G W O G R Y> 141
 C D E C I R Y P G C L H G T C Q Q P W O> 161
 C N C Q E G W G G L F C N Q D L N Y C T> 181
 H H K P C K N G A T C * T N T G Q G * 198
 S Y T * P S R * K N G G S L T D L * 213
 E N S Y S C T C P P G F Y G K I C E L S A M> 235
 T C A D G P G F N G G R C S D S P D G G> 255
 Y S C R C P V G Y S G F N C E K K I D Y> 275
 C S S S P C S N G A K C V D L G D A Y L> 295
 C R C Q A G F S G R H C D D N V D D C A> 315
 S S P C A N G G T C R D G V N D E S C T> 335
 C P P G Y T G R N C S A P A S R C E H A> 355
 P C H N G A T C H E R G H R Y * C E C A> 374
 R S Y G G P N C * F L L P E * P P G P * 391
 V V * L L L G C A A V V V C V R L R L Q K H> 412
 R P P A D P * R G E T E T M N N L * 428
 N C Q R E K D I S V S I I G * T O I K N T N> 449
 K K A D F H G D H * A D K N G F K A R Y P * 469
 V D Y N L V O D L K G D D T A V R D A H S K R D T K * 495
 Q P O G S S G E E K G T P * P T L R * G G * 514
 I * R K R P * S * S T * S K D * T * 526
 C V I * E V * 531

FIG. 14